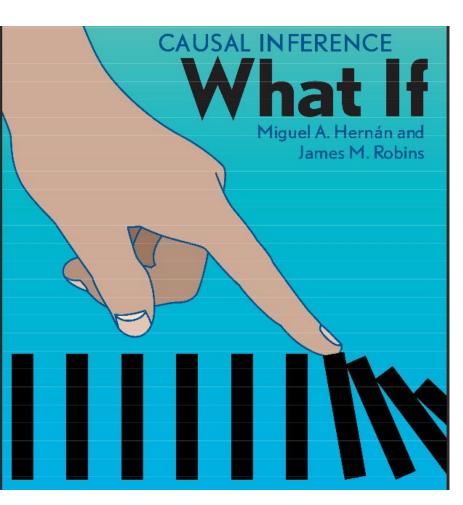
# book club: chapters 8.4 – 8.6



> selection bias & censoring

> adjustment for selection bias

> selection without bias

#### Selection bias review

```
n=1000
coin1 = rbinom(n,size=1,prob=0.5)
coin2 = rbinom(n,size=1,prob=0.5)

flips <- data.frame(coin1=coin1, coin2=coin2, selected= coin1 | coin2)
head(flips)</pre>
```

|        | coin1<br><int></int> | coin2<br><int></int> | selected<br>< g > |
|--------|----------------------|----------------------|-------------------|
| 1      | 0                    | 0                    | FALSE             |
| 2      | 1                    | 1                    | TRUE              |
| 3      | 1                    | 0                    | TRUE              |
| 4      | 1                    | 1                    | TRUE              |
| 5      | 0                    | 0                    | FALSE             |
| 6      | 0                    | 0                    | FALSE             |
| 6 rows |                      |                      |                   |

```
prob.coin= subset(flips,coin2==0)

paste0("p[coin1=head|coin2=tail]: ",signif(sum(prob.coin$coin1)/nrow(prob.coin),3))
```

```
[1] "p[coin1=head|coin2=tai1]: 0.498"
```



simultaneously flipping two coins







tail (0) coin1

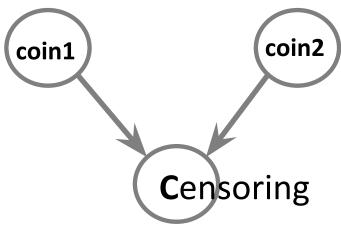
head (1) coin2

### Selection bias review

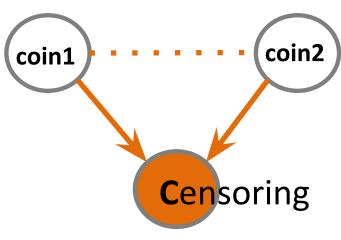
|      | coin1<br><int></int> | coin2<br><int></int> | selected<br>< g > |
|------|----------------------|----------------------|-------------------|
| 2    | 1                    | 1                    | TRUE              |
| 3    | 1                    | 0                    | TRUE              |
| 4    | 1                    | 1                    | TRUE              |
| 7    | 1                    | 1                    | TRUE              |
| 8    | 1                    | 1                    | TRUE              |
| 10   | 1                    | 1                    | TRUE              |
| rows |                      |                      |                   |
|      |                      |                      | Hid               |
|      |                      |                      |                   |

#### Selection bias review

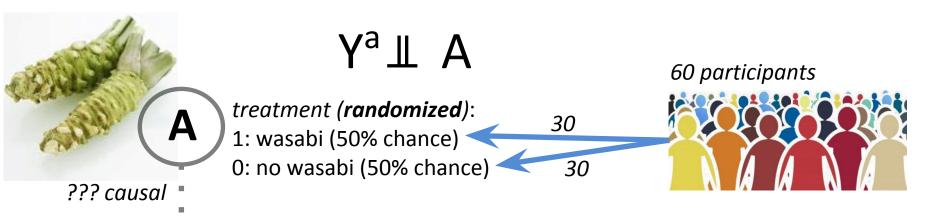


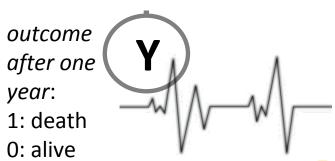


| Hide | prob.coin= subset(prob.coin,coin2==0) | paste0("p[coin1=head|coin2=tail]: ",signif(sum(prob.coin\$coin1)/nrow(prob.coin),3)) | | [1] "p[coin1=head|coin2=tail]: 1"



case I: we do get data from all participants



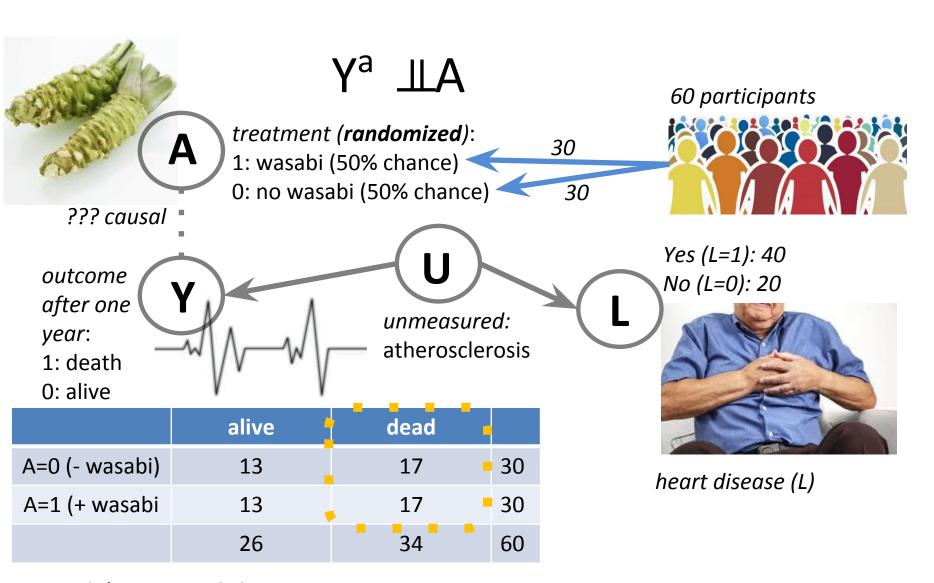


|                | alive | dead   |   |
|----------------|-------|--------|---|
| A=0 (- wasabi) | 13    | 17 - 3 | 0 |
| A=1 (+ wasabi  | 13    | 17 - 3 | 0 |
|                | 26    | 34 6   | 0 |

associational risk ratio:

$$Pr[Y^{a=1}=1] / Pr[Y^{a=0}=1] = 1$$

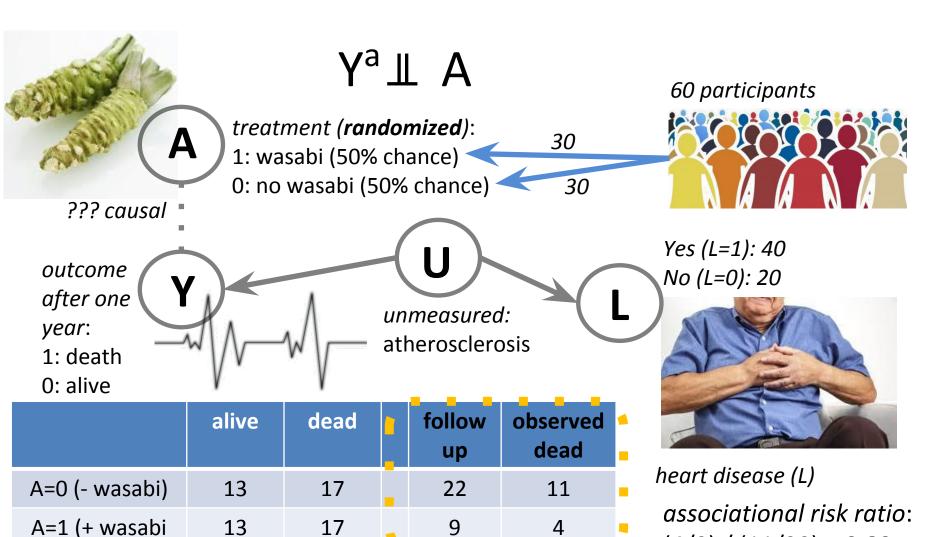
case I: we do get data from all participants



26

34

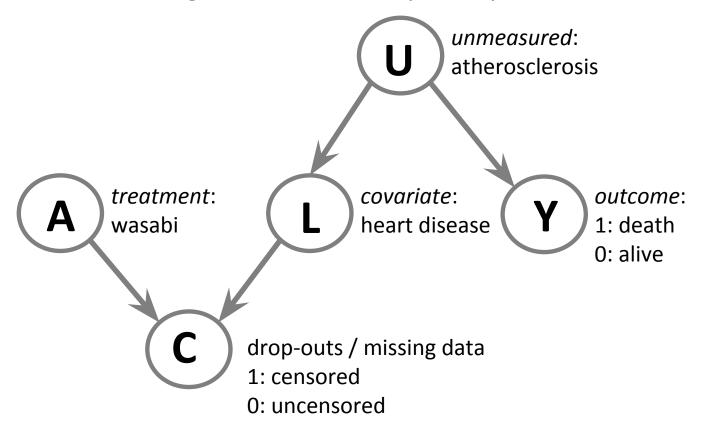
case II: we do NOT get data from all participants



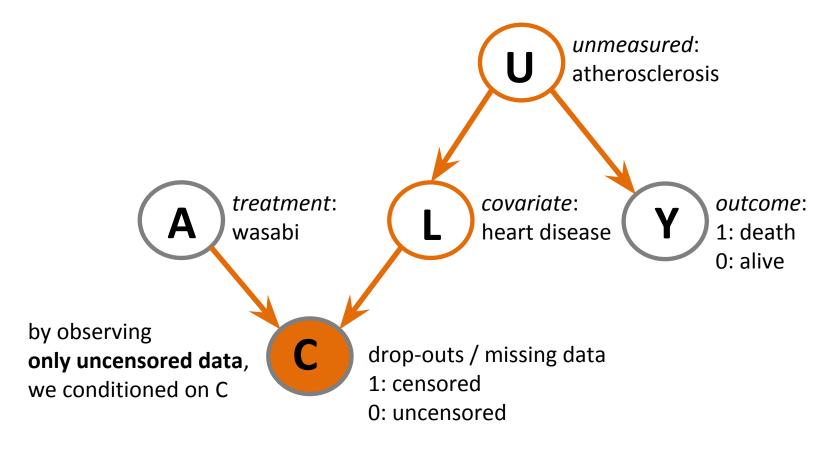
(4/9) / (11/20) =**0.89** 

on uncensored data

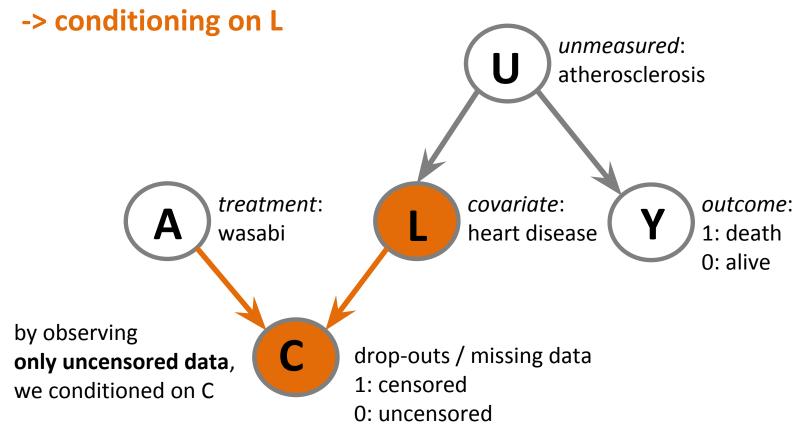
case II: we do NOT get data from all participants



case II: selection bias - conditioning on collider (backdoor path)



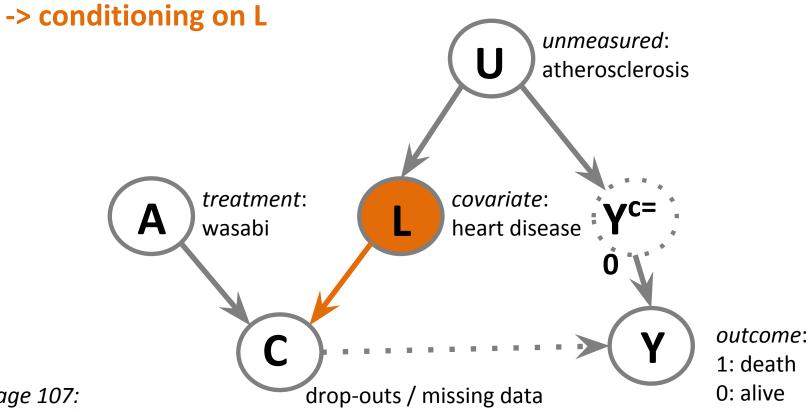
case III: selection bias - blocking the backdoor path



conditional risk ratio:

Pr[Y=1|A=1,C=0,L=I] / Pr[Y=1|A=0,C=0,L=I]

case III: selection bias - blocking the backdoor path



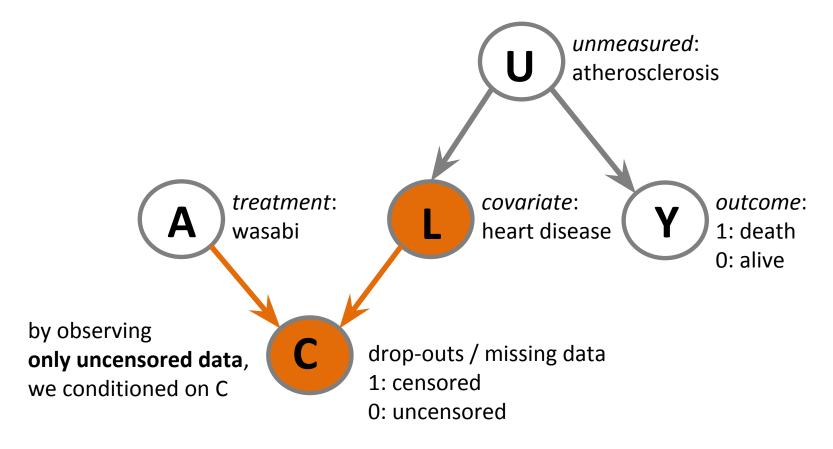
page 107:

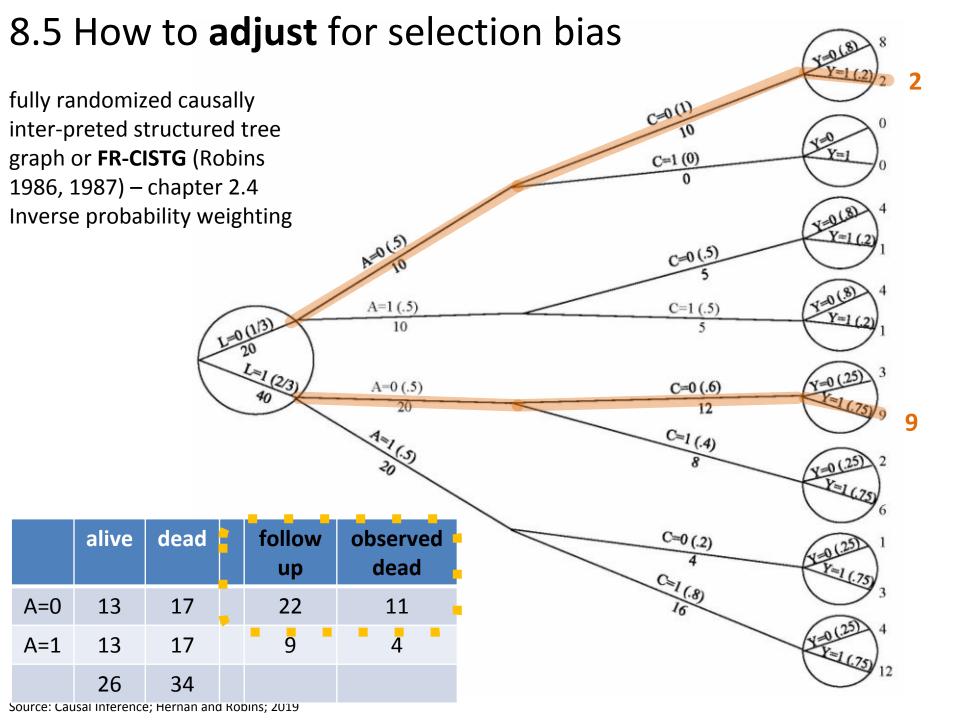
In causal diagrams with no arrow from censoring C to the observed outcome Y, we could replace Y by the counterfactual outcome  $Y^{c=0}$ and add arrows  $Y^{c=0} \longrightarrow Y$  and  $C \longrightarrow Y$ .

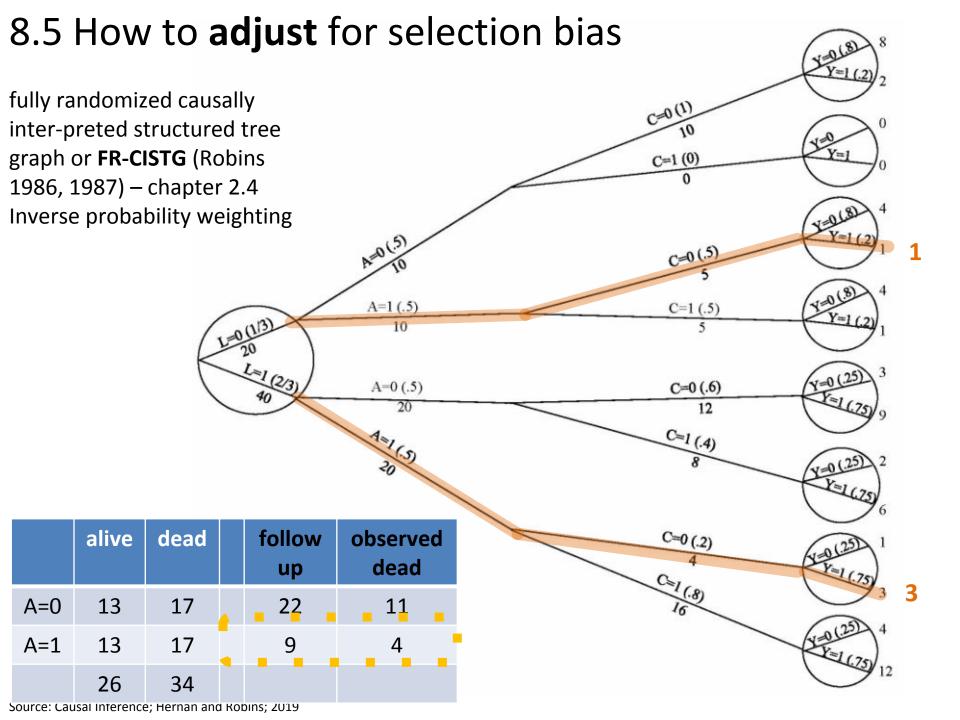
1: censored

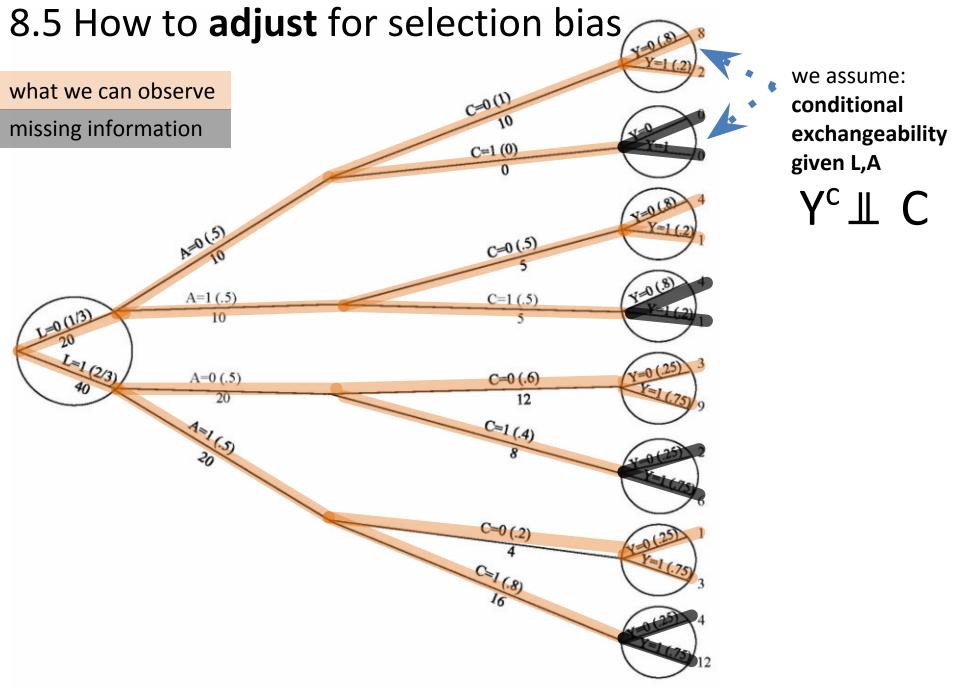
0: uncensored

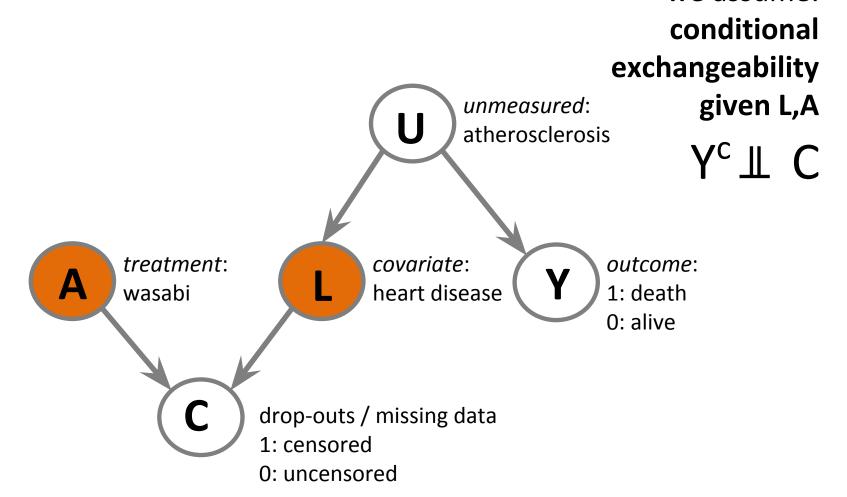
#### -> stratification











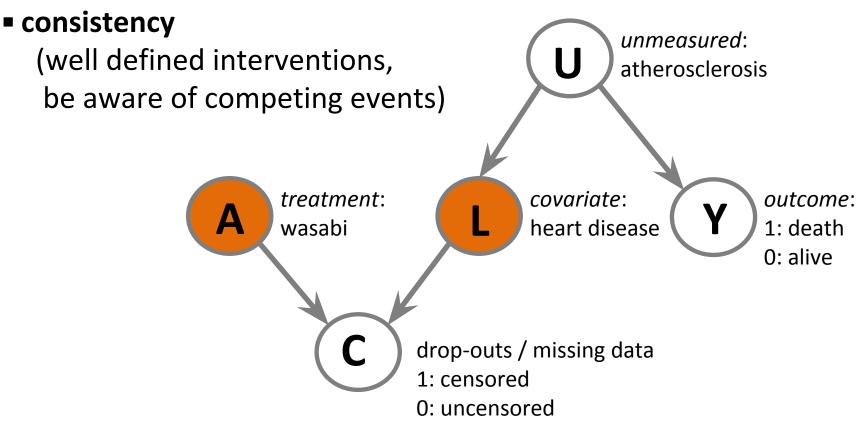
page 109:

conditioning on A and L is sufficient in blocking the backdoor path  $\mathbf{C} \leftarrow \mathbf{L} \leftarrow \mathbf{U} \rightarrow \mathbf{Y}$  between  $\mathbf{C}$  and  $\mathbf{Y}$ 

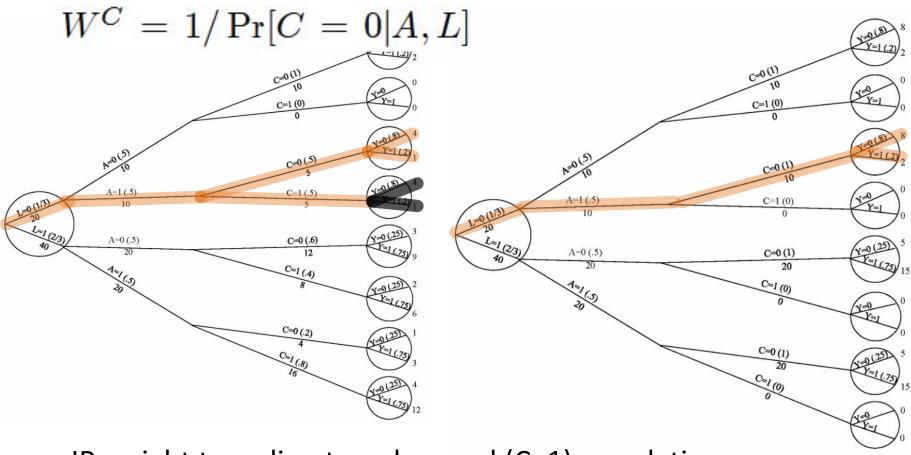
we assume:

identifiability conditions:

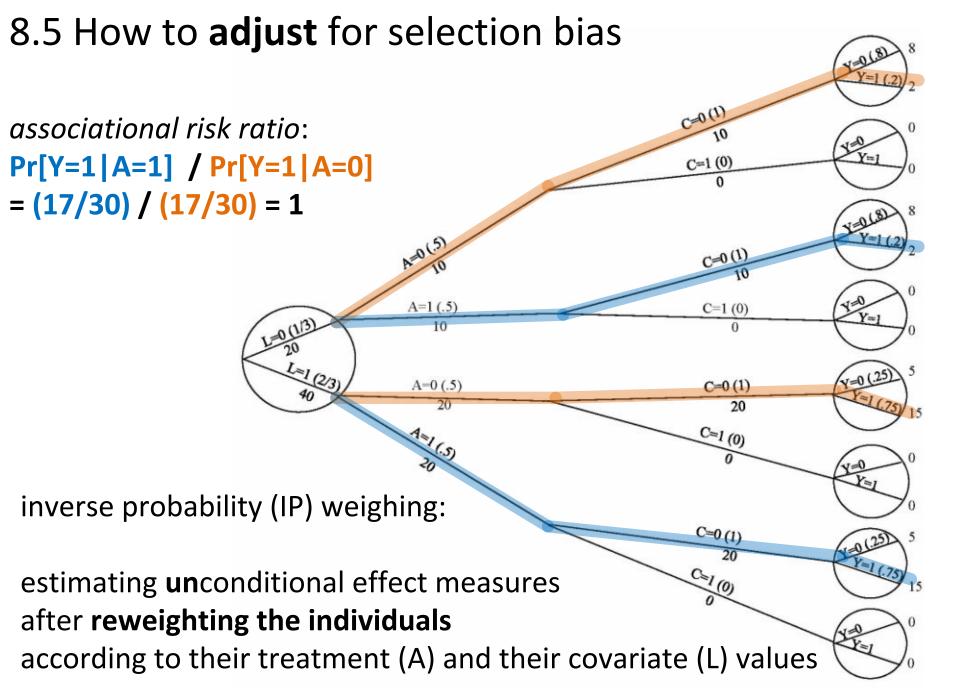
- exchangeability
- positivity

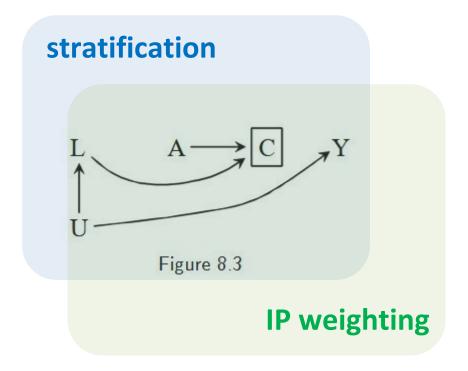


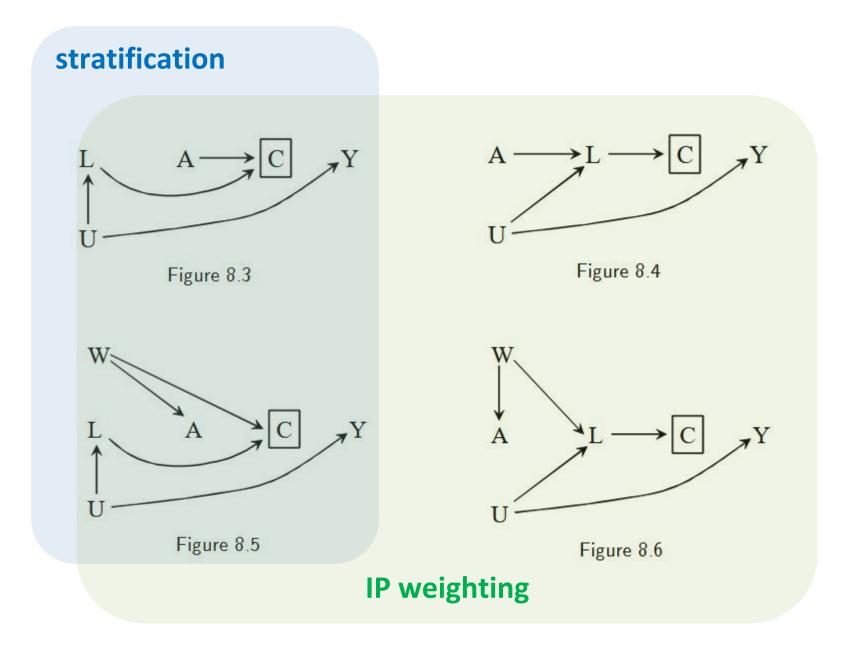
### -> inverse probability (IP) weighting



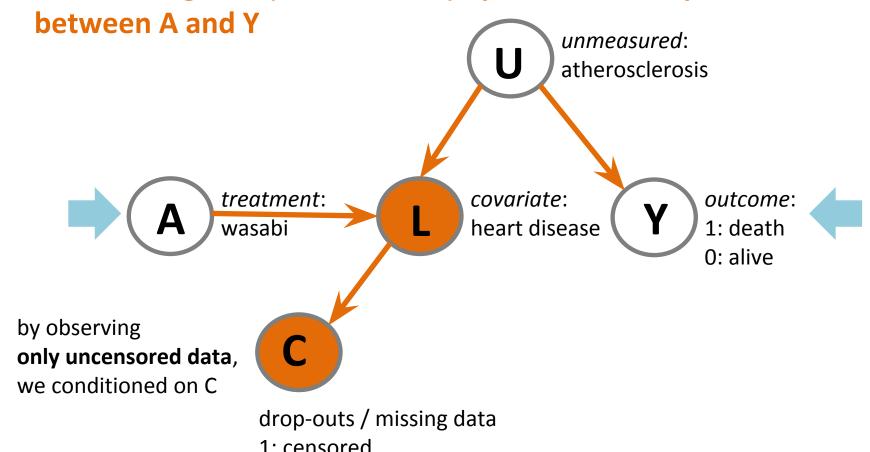
use IP weight to redirect unobserved (C=1) population to an estimated **pseudo-population** 







-> conditioning on L (stratification) opens backdoor path



1: censored

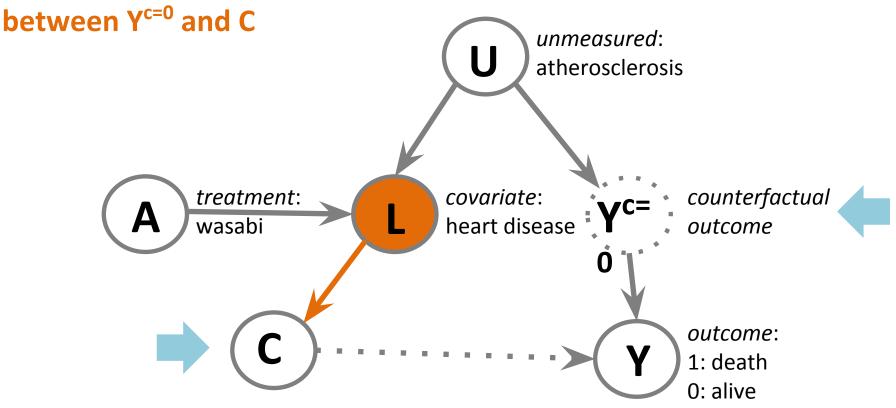
0: uncensored



**our focus** in verifying exchangeability

conditioning is represented by colored node

-> conditioning on L (IP weighting) d-seperates backdoor path

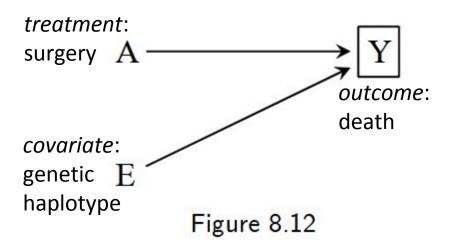




our focus in verifying exchangeability

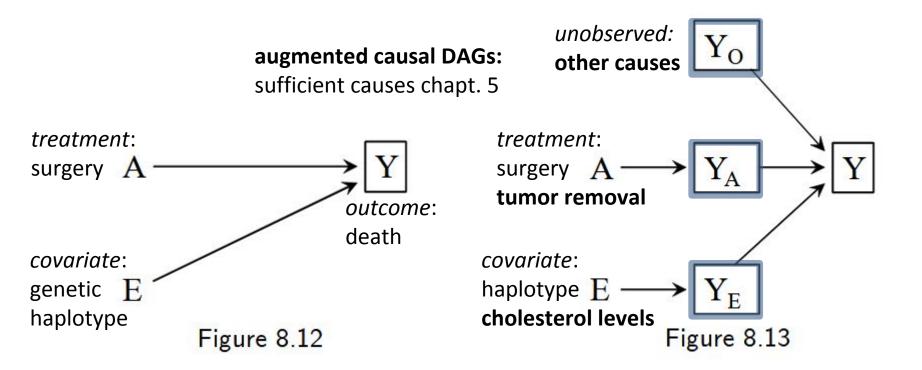
conditioning is represented by colored node

conditioning on the common effect Y induces conditional association between A and E



special situation (e.g. independent mechanisms):

dead Y=1: if 
$$(Y_0=1)$$
 or  $(Y_A=1)$  or  $(Y_E=1)$  alive Y=0: if  $(Y_0=0)$  and  $(Y_A=0)$  and  $(Y_E=0)$ 

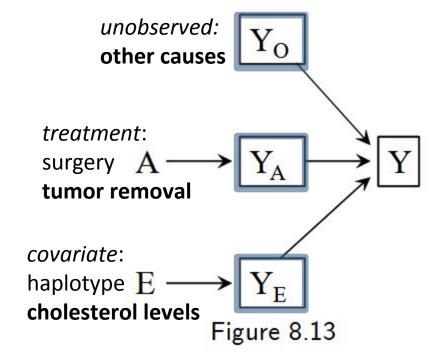


#### special situation (e.g. independent mechanisms):

A remains conditionally independent to E even if conditioning on the common effect Y within one stratum

dead Y=1: if 
$$(Y_0=1)$$
 or  $(Y_A=1)$  or  $(Y_E=1)$   
enforced determinism: alive Y=0: if  $(Y_0=0)$  and  $(Y_A=0)$  and  $(Y_E=0)$ 

the data in fig 8.13 follow a multiplicative survival model

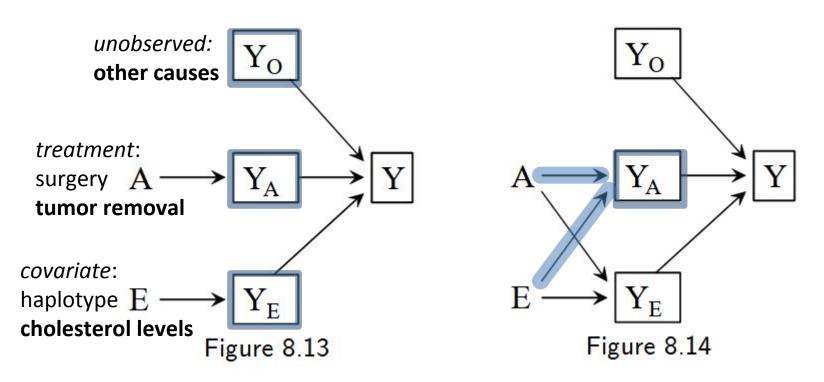


#### Technical Point 8.2

**Multiplicative survival model**. When the conditional probability of survival  $\Pr[Y=0|E=e,A=a]$  given A and E is equal to a product g(e)h(a) of functions of e and a, we say that a multiplicative survival model holds. A multiplicative survival model

$$\Pr\left[Y=0|E=e,A=a\right]=g(e)h(a)$$

is equivalent to a model that assumes the survival ratio  $\Pr\left[Y=0|E=e,A=a\right]/\Pr\left[Y=0|E=e,A=0\right]$  does not depend on e and is equal to h(a). The data follow a multiplicative survival model when there is no interaction between A and E on the multiplicative scale as depicted in Figure 8.13. If  $\Pr\left[Y=0|E=e,A=a\right]=g(e)h(a)$ , then  $\Pr\left[Y=1|E=e,A=a\right]=1-g(e)h(a)$  does not follow a multiplicative mortality model. Hence, when A and E are conditionally independent given Y=0, they will be conditionally dependent given Y=1.



#### common mechanisms:

conditioning on common effects renders A and E conditionally dependant

#### interpretation:

conditioning on a collider always induces an association beetween its causes, but this association could be restricted to certain levels of the common effect. Collider stratification is not always a source of selection bias.